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SEQUENCE LISTING

<110> Evotec NeuroSciences GmbH

<120> Diagnostic and therapeutic use of FOAP-13
polynucleotides and polypeptides for neurodegenerative
diseases

<130> 031985wo ME/BM

<140>

<141>

<150> 02019281.1

<151> 2002-08-28

<160> 18

<170> PatentIn Ver. 2.1

<210> 1

<211> 390

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: cDNA fragment
of the foap-13 gene

<400> 1

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cttggcctga ccctgggtctg gtctcagaat cacttttccc atctgtaaaa ttgagatgaa 180
ttttggtggt gaaagtctct cctggagcag atgtcctaga aggttttagg aatagtgaca 240
gagtcaggcc accccaaggg ccatgggagc cagctgacct gcttgaccga aggatttctg 300
acagactatc tttggggatg ttttcaagaa gggatataag ttatttactt tgggcattta 360
aaagaaaatt tctctcggga ataattttat                               390

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<210> 2

<211> 491

<212> PRT

<213> Homo sapiens

<400> 2

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Met Ala Gly Gln Gly Leu Pro Leu His Val Ala Thr Leu Leu Thr Gly
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Leu Leu Glu Cys Leu Gly Phe Ala Gly Val Leu Phe Gly Trp Pro Ser
      20              25              30

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Leu Val Phe Val Phe Lys Asn Glu Asp Tyr Phe Lys Asp Leu Cys Gly
      35              40              45

```

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Pro Asp Ala Gly Pro Ile Gly Asn Ala Thr Gly Gln Ala Asp Cys Lys
      50              55              60

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Ala Gln Asp Glu Arg Phe Ser Leu Ile Phe Thr Leu Gly Ser Phe Met
      65              70              75              80

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Asn Asn Phe Met Thr Phe Pro Thr Gly Tyr Ile Phe Asp Arg Phe Lys
      85              90              95

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Thr Thr Val Ala Arg Leu Ile Ala Ile Phe Phe Tyr Thr Thr Ala Thr
 100 105 110
 Leu Ile Ile Ala Phe Thr Ser Ala Gly Ser Ala Val Leu Leu Phe Leu
 115 120 125
 Ala Met Pro Met Leu Thr Ile Gly Gly Ile Leu Phe Leu Ile Thr Asn
 130 135 140
 Leu Gln Ile Gly Asn Leu Phe Gly Gln His Arg Ser Thr Ile Ile Thr
 145 150 155 160
 Leu Tyr Asn Gly Ala Phe Asp Ser Ser Ser Ala Val Phe Leu Ile Ile
 165 170 175
 Lys Leu Leu Tyr Glu Lys Gly Ile Ser Leu Arg Ala Ser Phe Ile Phe
 180 185 190
 Ile Ser Val Cys Ser Thr Trp His Val Ala Arg Thr Phe Leu Leu Met
 195 200 205
 Pro Arg Gly His Ile Pro Tyr Pro Leu Pro Pro Asn Tyr Ser Tyr Gly
 210 215 220
 Leu Cys Pro Gly Asn Gly Thr Thr Lys Glu Glu Lys Glu Thr Ala Glu
 225 230 235 240
 His Glu Asn Arg Glu Leu Gln Ser Lys Glu Phe Leu Ser Ala Lys Glu
 245 250 255
 Glu Thr Pro Gly Ala Gly Gln Lys Gln Glu Leu Arg Ser Phe Trp Ser
 260 265 270
 Tyr Ala Phe Ser Arg Arg Phe Ala Trp His Leu Val Trp Leu Ser Val
 275 280 285
 Ile Gln Leu Trp His Tyr Leu Phe Ile Gly Thr Leu Asn Ser Leu Leu
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 Thr Asn Met Ala Gly Gly Asp Met Ala Arg Val Ser Thr Tyr Thr Asn
 305 310 315 320
 Ala Phe Ala Phe Thr Gln Phe Gly Val Leu Cys Ala Pro Trp Asn Gly
 325 330 335
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 340 345 350
 Thr Gly Ser Ser Thr Leu Ala Val Ala Leu Cys Ser Thr Val Pro Ser
 355 360 365
 Leu Ala Leu Thr Ser Leu Leu Cys Leu Gly Phe Ala Leu Cys Ala Ser
 370 375 380
 Val Pro Ile Leu Pro Leu Gln Tyr Leu Thr Phe Ile Leu Gln Val Ile
 385 390 395 400
 Ser Arg Ser Phe Leu Tyr Gly Ser Asn Ala Ala Phe Leu Thr Leu Ala
 405 410 415
 Phe Pro Ser Glu His Phe Gly Lys Leu Phe Gly Leu Val Met Ala Leu

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420	425	430
Ser Ala Val Val Ser Leu Leu Gln Phe Pro Ile Phe Thr Leu Ile Lys		
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Gly Ser Leu Gln Asn Asp Pro Phe Tyr Val Asn Val Met Phe Met Leu		
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Ala Ile Leu Leu Thr Phe Phe His Pro Phe Leu Val Tyr Arg Glu Cys		
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Arg Thr Trp Lys Glu Ser Pro Ser Ala Ile Ala		
485	490	

<210> 3

<211> 2630

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: cDNA of the human foap-13 gene

<400> 3

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tgccacgggt taaattttca ggtgaagagt gaggttgtca tggcctcagc tatgcttctc 2220

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ggctctccct caagagtga gccttggcta gagaactcac agctctggga aaaagaggag 2280
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<210> 4

<211> 13

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: one-base
anchor oligonucleotide

<400> 4

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13

<210> 5

<211> 13

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: one-base
anchor oligonucleotide

<400> 5

httttttttt ttg

13

<210> 6

<211> 13

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: one-base
anchor oligonucleotide

<400> 6

httttttttt ttc

13

<210> 7

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for the
foap-13 gene

<400> 7

tcaggtgaag agtgaggttg tca

23

<210> 8

- 5 -

<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: primer for the
foap-13 gene

<400> 8
ggctgcactc ttgagggaga 20

<210> 9
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: primer for the
cyclophilin B gene

<400> 9
actgaagcac tacgggcctg 20

<210> 10
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: primer for the
cyclophilin B gene

<400> 10
agccgttggt gtctttgcc 19

<210> 11
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: primer for the
ribosomal protein S9

<400> 11
ggtcaaattt accctggcca 20

<210> 12
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: primer for the
ribosomal protein S9

<400> 12
tctcatcaag cgtcagcagt tc 22

<210> 13
<211> 19
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for the
beta-actin gene

<400> 13
tggaacggtg aaggtgaca

19

<210> 14
<211> 19
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for the
beta-actin gene

<400> 14
ggcaaggac ttcctgtaa

19

<210> 15
<211> 20
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for the
GAPDH gene

<400> 15
cgtcatgggt gtgaaccatg

20

<210> 16
<211> 21
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for the
GAPDH gene

<400> 16
gctaagcagt tgggtgtgca g

21

<210> 17
<211> 21
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for the
transferrin receptor (TRR)

- 7 -

<400> 17
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21

<210> 18
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: primer for the
transferrin receptor (TRR)

<400> 18
agcagttggc tgttgtacct ctc

23